SEQUENCE LISTING

<110> Steinkasserer, Alexander
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Ala Pro Ala Thr Pro Glu Val Lys Val Ala Cys Ser Glu Asp Val Asp 20 25 30
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Leu Pro Cys Thr Ala Pro Trp Asp Pro Gln Val Pro Tyr Thr Val Ser
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Trp Val Lys Leu Glu Gly Gly Glu Glu Arg Met Glu Thr Pro Gln 50 55 60
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Thr Thr Ser Cys Asn Ser Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro 100 105 110
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Cys Pro Ala Gln Arg Lys Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu 130 135 140
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	٠.		100		Ser			105					110			
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gaa Glu	agg Arg	ccc Pro	tat Tyr 75	tcc Ser	ctg Leu	aag Lys	atc Ile	cga Arg 80	aac Asn	act Thr	acc Thr	agc Ser	tgc Cys 85	aac Asn	tcg Ser	288
GJ À aaa	aca Thr	tac Tyr 90	$\operatorname{\mathtt{Arg}}$	tgc Cys	act Thr	ctg Leu	cag Gln 95	gac Asp	ccg Pro	gat Asp	GJÀ āāā	cag Gln 100	aga Arg	aac Asn	cta Leu	336
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Gly .40	Gly	Glu	Glu	Arg	Met 45	Glu	Thr	Pro	Gln	Glu 50	Asp	His	Leu	Arg	Gly 55	

Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn 65 Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser 80 Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu 95 Ser Gly Lys Val Ile Leu Arg Val Thr Gly Cys Pro Ala Gln Arg Lys Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile 125 <210> 9 <211> 435 <212> DNA <213> Artificial Sequence <220> <2235 Description of Artificial Sequence: partial sequence of pGEX2ThCD83ext_mut129_CtoS <221> CDS <222> (1)..(417) <220> <221> mat_peptide <222> (28)..(417) cct cca aaa tcg gat ctg gtt ccg cgt gga tcc ccg gga acg ccg gag 48 Pro Pro Lys Ser Asp Leu Val Pro Arg Gly Ser Pro Gly Thr Pro Glu gtg aag gtg gct tgc tcc gaa gat gtg gac ttg ccc tgc acc gcc ccc 96 Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro tgg gat ccg cag gtt ccc tac acg gtc tcc tgg gtc aag tta ttg gag 144 Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu 30 ggt ggt gaa gag agg atg gag aca ccc cag gaa gac cac ctc agg gga 192 Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly 45 cag cac tat cat cag aag ggg caa aat ggt tct ttc gac gcc ccc aat 240 Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn gaa agg ccc tat tcc ctg aag atc cga aac act acc agc tgc aac tcg 288 Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser ggg aca tac agg tgc act ctg cag gac ccg gat ggg cag aga aac cta 336 Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu 95 agt ggc aag gtg atc ttg aga gtg aca gga tcc cct gca cag cgt aaa Ser Gly Lys Val Ile Leu Arg Val Thr Gly Ser Pro Ala Gln Arg Lys

<213> Artificial Sequence

<223> Description of Artificial Sequence: primer

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antisense-CD83extra mu	tant	:C	tc	ıS
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